

## Kernelized SVM for classification

Kernel trick: replacing the linear transformation that we usually have in linear models with a kernel - any function that measures the distance between two vectors.

Linear models defined in terms of the kernel function:

$$y(x; \alpha) = \sum_{n=1}^N \alpha_n k(x_n, x)$$

$$y(x; \alpha) = \sum_{n=1}^N \alpha_n x_n^T x$$

Solution

$$\alpha = (K + \lambda I_N)^{-1} t$$

linear model with linear kernel

Gram matrix

$$K = \begin{bmatrix} k(x_1, x_1) & \dots & k(x_1, x_N) \\ \vdots & \ddots & \vdots \\ k(x_N, x_1) & \dots & k(x_N, x_N) \end{bmatrix}$$

The evaluation of  $\alpha$  would be almost infeasible but an important property of SVMs is that the computation of the model parameters corresponds to a convex optimization problem and so any local solution is also a global optimum.

Previous formulation of the SVMs:

$$\left. \begin{aligned} y(x; w) &= w^T x + w_0 \\ \hat{w} &= \sum_{n=1}^N \alpha_n^* t_n x_n \end{aligned} \right\} y(x) = w_0 + \sum_{x_n \in \text{SV}} \alpha_n^* t_n x_n^T x_n$$

Hyperplanes expressed with support vectors

$$y(x) = w_0 + \sum_{x_i \in \text{SV}} \boxed{\alpha_i^* t_i} x_i^T x_i = 0$$

Samples on one side are classified one way, samples on the other side are classified the other way

$$y(x'; \alpha) = \text{sign} \left( w_0^* + \sum_{x_k \in \text{SV}} \boxed{\alpha_k^* t_k} x_k^T x' \right)$$

Solution of the problem:

$$w^* = \sum_{n=1}^N \boxed{\alpha_n^* t_n} x_n$$

linear kernel

Now with kernelization:

$$x_i^T x_j \longleftarrow k(x_i, x_j)$$

Classification:

$$y(x'; \alpha) = \text{sign} \left( w_0 + \sum_{n=1}^N \alpha_n \boxed{k(x_n, x)} \right)$$

generic kernel